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[F.]F) using at least one extended [immobilised] immobilized nucleic acid strand to repeat steps D) and E), so as to provide additional extended [immobilised] immobilized nucleic acid strands and, optionally, [G.]G) repeating step F) one or more times.

- (Twice Amended) A method according to claim 1, wherein said single-3. stranded target nucleic acid [is produced by providing] comprises a given nucleic acid sequence to be amplified (which sequence may be known or unknown) [and adding thereto] to which have been added a first nucleic acid sequence and a second nucleic acid sequence; wherein said first nucleic acid sequence [hybridises] hybridizes to one of said plurality of primers and said second nucleic acid sequence is complementary to a sequence which [hybridises] hybridizes to one of said plurality of primers.
- (Twice Amended) A method according to claim 1; wherein said single-4. stranded target nucleic acid [is produced by providing] comprises a given nucleic acid sequence to be amplified (which sequence may be known or unknown) [and adding thereto] to which have been added a first nucleic acid sequence and a second nucleic acid sequence;. wherein said first nucleic acid sequence [hybridises] hybridizes to one of said plurality of primers and said second nucleic acid sequence is the same as the sequence of one of said plurality of primers.
- (Twice Amended) A method according to claim 3 wherein said first and 5. second nucleic acid sequences are provided at [first and second] 3' and 5' ends of said singlestranded target nucleic acid.
- (Previously Amended) A method according to claim 3, wherein a tag is also added to 6. the given nucleic acid sequence, said tag enabling amplification products of the given nucleic acid sequence to be identified.
- (Previously Amended) A method according to claim 1 wherein the plurality of primers 7. is a plurality of primers that have the same sequence.
- (Previously Amended) A method according to claim 1, wherein the plurality of primers comprises at least two different types of primer, one type having a different sequence from another type.

		Eric Kawashima et a Application No.: 09/402,277 Page 8	
•	1 2	9. (As filed) A method according to claim 8, wherein the plurality of primers consists of 2 ⁿ different types of primer; wherein n is an integer.	
	1	10. (As filed) A method according to claim 9, where n is 2.	
	1 2	11. (Twice Amended) A method according to claim 8, wherein the different types of primer are present in [substantially] about the same concentrations as one another.	
	1	12. (Twice Amended) A method according to claim 1, wherein the primers are [substantially] homogeneously dispersed over a given area.	
\wedge	2 1 SuDE	13. (Twice Amended) A method according to claim 1, wherein the primers	
	$\frac{2}{1}$	are located in a predetermined arrangement (e.g. in a grid pattern). 14. (Twice Amended) A method according to claim 1, wherein a supply of	
	2	nucleotides and a nucleic acid polymerase are used to extend primers.	
	1 2	15. (Twice Amended) A method according to claim 1, wherein heating is used to separate annealed nucleic acid strands.	_
	1 2	16. (Previously Amended) A method according to claim 14, wherein the nucleic acid polymerase is not rendered inactive by the heating conditions used to separate annealed nucleic acid strands.	-
	1 2 3 3 2 2 3	polymerase is taq polymerase, or is another polymerase that is derivable from a thermophilic organism; or is a thermostable derivative thereof. 18. (Twice Amended) A method according to claim 1, wherein said primer extension results in the incorporation of one or more detectable labels (e.g. fluorescent labels or radiolabels) into extended [immobilised] immobilized nucleic acid strands. 19. (Twice Amended) A method according to claim 1, further including the step of treating one or more extended [immobilised] immobilized nucleic acid strands so as to release a nucleic acid molecule or a part thereof.	
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Eric Kawashima et a Application No.: 09/402,277 Page 9

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(As Filed) A method/according to claim 19, wherein said treating consists of cleavage 20. with a restriction endonuclease or with a ribozyme.

- (Previously Amended) A method according to claim 1, wherein one or more of said 21. primers has a restriction endonuclease recognition site or a ribozyme recognition site or has part of such a site, which part becomes complete when primer extension occurs.
- (Previously Amended) A method according to claim 1 that is automated to allow 22. repeated cycles of nucleic acid amplification.
- (Previously Amended) A method according to claim 1, when used to amplify a 23. plurality of different nucleic acid sequences
- (As Filed) A/method according to claim 23, when used to amplify a plurality of 24. different nucleic acid sequences simultaneously.
- (Previously Amended) A method according to claim 23, wherein said different nucleic 25. acid sequences are each provided with a first and second nucleic acid sequence as described in any of claims 3 to 5, said first and second nucleic acid sequences being the same for the each of the different nucleic acid sequences.
- (Previously Amended) A method according to claim 23, wherein said different nucleic 26. acid sequences are each provided with a different tag so that the different sequences can be distinguished from one another.
- (Amended) A plurality of [immobilised] immobilized nucleic acids 27.. producable by a method according to any preceding claim.
- (Amended) A plurality of [immobilised] immobilized nucleic acids in 28. the form of one or more distinct areas on a surface, each area comprising a plurality of identical nucleic acid strands and a plurality of dentical complementary strands thereto; wherein each nucleic acid strand within such ar area is located so that another nucleic acid strand is located on the surface within a distance of the length of that strand.
- (Amended) A plurality of [immobilised] immobilized nucleic acids according to claim 27 or claim 28, wherein there is at least one distinct area present per mm² of surface on which the nucleic acids are [immobilised] immobilized.

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Eric Kawashima et a Application No.: 09/402,277 Page 10 (Amended) A plurality of [immobilised] immobilized nucleic acids 30. according to claim 28, wherein the number of distinct areas/mm² of surface on which the nucleic acids are [immobilized is greater than 1, greater than 10^2 , greater than 10^3 or greater than 10^4 . (Twice Amended) An apparatus for performing a method as described 59. 1 in claim 1; comprising a plurality of [immobilised] immobilized primers, a nucleic acid 2 polymerase, a plurality of nucleotides and means for separating annealed nucleic acid strands. 3 (As Filed) An apparatus according to claim 59, wherein the means for separating 60. 1 annealed nucleic acid strands comprises a controlled heating means. (Twice Amended) An apparatus for [analysing] analyzing a plurality of 61. nucleic acid molecules [according to claim 27] producable by a method according to claim 1, 2 wherein said apparatus comprises a source of reactants and detector means for detecting one or 3 more signals produced after said reactants have been applied to said nucleic acid molecules. 4 (Previously Amended) An apparatus according to claim 61 wherein said detector 1 means has sufficient resolution to distinguish between the distinct areas on a surface, each area comprising a 2 plurality of identical nucleic acid strapds and a plurality of identical complementary strands thereto; wherein each 3 nucleic acid strand within such an area is located so that another nucleic acid strand is located on the surface 4 within a distance of the length of that strand. 5 (Previously Amended) An apparatus according to claim 61 comprising a charge 63. 1 coupled device (CCD). 2 (As Filed) An apparatus according to claim 63 wherein said charge coupled device 64. 1 (CCD) is operatively connected with a magnifying device (e.g. a microscope). 2 (Twice Amended) A kit for use in screening, diagnosis or in nucleic 65. 1 acid sequencing; comprising a plurality of [immobilised] immobilized nucleic acid according 2 3 to claim 27.